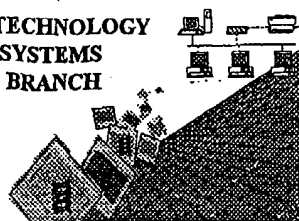# RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: _09/404,520A_

Source: _1,600_

Date Processed by STIC: _1/7/2004_

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221
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   http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
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Revised 10/08/03

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1600

RAW SEQUENCE LISTING   DATE: 01/07/2004
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Input Set : E:\ENU.corrected SEQ list 12.23.03.rpt.txt
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pp 1-16

Does Not Comply
Non-Use or Defective Needed

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 2 <110> APPLICANT: Cao, Yongwei
 3         Ghodssi, Azita
 4         Hinkle, Gregory J.
 5         McIninch, James
 6         Timberlake, William E.
 7         Yu, Jaehyuk
 9 <120> TITLE OF INVENTION: Emericella nidulans Genome Sequence and Uses Thereof
11 <130> FILE REFERENCE: 38-10(15498)A
13 <140> CURRENT APPLICATION NUMBER: US/09/404,520A
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ERRORED SEQUENCES

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RAW SEQUENCE LISTING   DATE: 01/07/2004
PATENT APPLICATION: US/09/404,520A   TIME: 12:07:42

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10795 taagcctatc acgttaacac ctcttaccaa cctgtcactt cggccaagtc cactctccat  120
10797 ccgtttcttg cgccgttgaa gcttgaagag ggtgatggct gcacgcagca ctgtctttat  180
10799 acctggccat tgcatttgtc tctttactat ccaaaatttc ctgcgcgtat ctcccagaat  240
10801 tacaggccct gcttctaaac agatagtctc agccgtgaca gctggtccta tccgttgtct  300
10803 gagaaatacg ggccagggtg ccctccgccg attttatatg gcaatccaag cactaaacga  360
10805 ttcgatgcaa ttgatgaaac tcctcgcaga gcctacacct caacgtcgag gtgatataac  420
10807 tgcccgtgca ggtagcgcat tagggctttg tacatgctac gcgcaacgca aaactcaaat  480
10809 catgtttta cgcctgagac catcggtata gtgttatctt ctaaagggac gtacctacca  540
10811 gcgtttctga caactatttt attatagcgg taactttgca gatatagtct ttatgaaggt  600
10813 tttcgcaaag tatgagccgt caatttcgcc aggatatctt gggccactgg gggtgagatt  660
10815 gttggccacg tgatataaat ggcctgatct ccaaccccctg acgaactgac ccctcaatat  720
10817 accgtattga gagacacaga agaagagatt tatcgccaac gagatagata gtcaatcgtc  780
10819 atatggcatc cgctccttag caagctacat ccgtgtagtt gatacagttt ccttctgccc  840
10821 tttcccctttt gagcgtacat gcgtatctac aacaatgcga ttattaccaa aaacaagtca  900
10823 ggtccaaata ggccttaaaa cgccgcagtt catctaattg agcaatggaa ctaggggtcc  960
10825 tcgaagcggg ctcggctgct cccgtaacgc ctgagtccaa tccccgcccg ccccccgctca 1020
10827 aatcactcct ataactctca aaattttcca cgtacctggg aatctctcag attttttccaa 1080
10829 ttggcgctcc gcccaccgct cacttttcct tccaggcgaa aagacagcta gcatgtcatc 1140
10831 tgaaaccatc cctcagaagc agaagcaccc gcaaggacag ggcgccaaga gctcccgatg 1200
10833 gcgtaacgcg cttggcaggg actgggcttt ggtatccctt ttcaccggct taatgaccgc 1260
10835 agtaatcggt tggtggcttg gaatcctgaa tgcagaaatg gcgttggaag cgccttttgg 1320
10837 attttttccag cgtatcgata tagacagcca ttgggaggag caccgaaaag aggtgaaaga 1380
W--> 10839 cactttcatg gccagctggc acgcctacgc caaacatgcc tggggtaaga ngatcatcag 1440
10841 cttcaaattc ggcttctttt ctatttttat tattattttg tgttt                   1485
11004 <210> SEQ ID NO: 149
11005 <211> LENGTH: 608
11006 <212> TYPE: DNA
11007 <213> ORGANISM: Aspergillus nidulans        insert <2207
11009 <223> OTHER INFORMATION: unsure at all n locations
E--> 11010 <400> SEQUENCE: 149
11012 ctagaactaa gaactaggga tcaattacag gaagattcgg acaaagaata gatggctaac   60
11014 tacgatctat gccaactatt tcgaggctaa aggagtgtaa tcaaagaaac gaataggtgc  120
11016 aactatggct gctatggccc caggggtggg aagaatcagg caagccctag atcacctgca  180
```

RAW SEQUENCE LISTING   DATE: 01/07/2004
PATENT APPLICATION: US/09/404,520A   TIME: 12:07:42

Input Set : E:\ENU.corrected SEQ list 12.23.03.rpt.txt
Output Set: N:\CRF4\01072004\I404520A.raw

```
        11018 cgtccaccca gaagcgggtg tcgtcagact gatatttcga atcactctca atctgctgct  240
        11020 gagtccatgc catagcgtcg atgtatcctc tagaagaata tcagtgaaga ccccccaaaa  300
W-->    11022 tcacaatcgt tggatggcga gatactcgcc cttgatgtcc tcggtgaggg ngattgttca  360
        11024 caccagactc gcatttgcgg tggccgctga accagatctc atatcaaaat ggttttcacc  420
        11026 ttcgtagcga gcttgaatat ccttggccaa gtcggagctt gcatccatcg ggtcaataga  480
        11028 gagacagagg ccggcgagct tgacccggaa gaagagccct tatttgagaa tggagccggt  540
        11030 atcgccctgg tcgatgggcc tgcctagaca ctttgacgcc caatgaccca gaatgaacaa  600
        11032 ttctggtt                                                            608
        11279 <210> SEQ ID NO: 154
        11280 <211> LENGTH: 2081
        11281 <212> TYPE: DNA
        11282 <213> ORGANISM: Aspergillus nidulans    insert <220?
        11284 <223> OTHER INFORMATION: unsure at all n locations
E-->    11285 <400> SEQUENCE: 154
        11287 tgcctcggat actcgaaaga gtaacatgtc ggtgtgcagc agcggcgaga gaacccaacc   60
        11289 gttcctcttt tgatacatgt ctcccagttg aggatacagg taaccaggta atcaacaact  120
        11291 tggatgatac tgcctgatgg cctcgtgaag ctcgacaggg ccgccgttgg acataaaaag  180
        11293 atcaatatcc cgcagttgag ggatcttatg aatgaggtca cttaccctgg cgttgaggtt  240
        11295 cgttgaacaa ccctcgactt tgggagtgct tcgaactccc aagttctcag cttgaagcgc  300
        11297 agaccggagg ccgctgtagg cacgaagcac tggtgcgcgt cgtgggccgc taatcggata  360
        11299 ccatgggacg atgatactga gtcgtttttgc gtgaacgagc acacggctgg cttctggctg  420
        11301 ccggagaaac ttaagaggca ttatccttga gtctacgtat gatgatatgc tcgtacaaga  480
        11303 gcgacctggt aaccctgttt cactcttggt ccacctcatc acgtatctgc agacacggta  540
        11305 agatataccA ccattgaaaa agggttaacg tacttgtctc ggaatcagac atttaaactc  600
        11307 tgatggcagt ccgtccatgt tttggtagag gggttggatg gtgtgcggag ggtttttaaaa  660
        11309 gaaatgctag cttggggaac atggctttct tttataatca ttctacccag agcttaccta  720
        11311 acagcacaga atagcctact tcctgcctag cggaaaacat atctagacct agatagatct  780
        11313 cctgctctga tcctccccct ctagccgcac ctaaacccaa caacaccta gacatcagtc  840
        11315 ttcccacact aattcttgta gaaactgacg atctcggtgt tccccttttc cgcatcgtac  900
        11317 ggtgggagcc cgtaaacgtg ccagtttttc tcactgcaga gtggcttcct ccaatttggc  960
        11319 aggtagttat tgaggcagct cagcctcccg aaattgactt tatcctttga gagtatctgt 1020
        11321 tatgaatgct caaaggggaa agggtagaag gaagctgcat caactacacg aacgtagcct 1080
        11323 actaaggata ggatgccata agacgattga ttctctatct tgatgacgat agttctcact 1140
        11325 tcttcaatgt acaactcctc gagggatcag gccaggggcc agagatcatg ccatacagat 1200
        11327 cacgtggcac gtgacacgag gctcaacatg atatattcgt atcagtatcc tatgtgataa 1260
        11329 gtctgttctt cctcgtccaa aaatcaccaa cagtattaat cgataaatga tgtgctcggt 1320
        11331 agtatgttag atatataacc tcacttctat aactatagat ttgggataca ggtacagaac 1380
        11333 catgtcttgc tatgtaggtt cataagccag tgtcataact taattcaatt ttcttgccaa 1440
        11335 gagactgaca aacattgtgt atcattgcgt cttttaagta cggcatactg ttcgaacaaa 1500
        11337 agggcaaaaa cgtatttaac acgtaacgag tttgtaccag ctcagattct gccagcagaa 1560
        11339 atttgattac atatgcagga tagcttgctt agtttcaccg actggttact catcctaagc 1620
        11341 accacgtaaa ggtgcaagct atgctcaggt gaggtacaca gcccgtaccc ctaggcagtg 1680
        11343 ctggtgttcg tacaggaaaa tcaagaaagc tcgtacggac cttataagac aatcggattg 1740
        11345 tcctcatccc cttcagggaa atcaaggtag ataggatttt cagcctcacc ttccactgga 1800
        11347 tcaagattga taaccagctg gtttgcagcc ccctgggta gaccaagccc atcaccaagc 1860
        11349 ataccctgcc agtgctcgtc tcgtacggag cagcaatgcc catggtcacg ccatttacaa 1920
        11351 ttggcgcagc atccgttgaa tgccccaggc acatggcggc attcagggaa gactggtctt 1980
        11353 tcttgacgga gcggcatcct ttacagggct ggtcagacga gtcccccaga ctgtatagca 2040
W-->    11355 ttgcattgat gtcnangggc cggtgttgca taattgatgc a                       2081
```

RAW SEQUENCE LISTING   DATE: 01/07/2004
PATENT APPLICATION: US/09/404,520A   TIME: 12:07:42

Input Set : E:\ENU.corrected SEQ list 12.23.03.rpt.txt
Output Set: N:\CRF4\01072004\I404520A.raw

```
    11958 <210> SEQ ID NO: 166
    11959 <211> LENGTH: 173
    11960 <212> TYPE: DNA
    11961 <213> ORGANISM: Aspergillus nidulans      insert <2207
    11963 <223> OTHER INFORMATION: unsure at all n locations
E--> 11964 <400> SEQUENCE: 166
    11966 taaagggatc tagaatcgtg attatgagtt tgataaggga aagcacatgg aatacattgg   60
W--> 11968 aaattagtgc tataaggcat cgctangaca gaaggcagaa tactgtgttg tataggttcc  120
    11970 gagagaatga ctgttgagtc gatgtgttca ttagtggttt ccattttagt cag          173
    13005 <210> SEQ ID NO: 177
    13006 <211> LENGTH: 712
    13007 <212> TYPE: DNA
    13008 <213> ORGANISM: Aspergillus nidulans   insert <2207
    13010 <223> OTHER INFORMATION: unsure at all n locations
E--> 13011 <400> SEQUENCE: 177
    13013 cctcggacaa caaggtctac ttccagctgc accagaacgg ctcctcgaca gttaacacgg   60
    13015 gccccaatgg attggagcgt ctcgactaca tcatctctgg ggccgagaag cgcggaatca  120
    13017 agttggtcat cccgttggtt aactactggg acgactttgg tggcatgaat gcctacatca  180
    13019 gcgcctatgg cggcgacaag cccggctggt acaccaatga taagattcag gctgcatatc  240
    13021 acgcctacgt caaggccgtg gtgagccgct atgttgactc ccggcaatct ttgcctggga  300
    13023 gttggcaaat gagccccgct gcagtggttg cgacacgtcc atcatcaatc agtgggcgac  360
    13025 gaaaaccagc agcttcatta aatccctgga tccaaaccat atggtagcca tgggcgacgg  420
    13027 tgagtccct gaatctccct ctctctggtc gtagaccgcc gctaagaggg gatctctctt  480
    13029 agagggcatg ggcctccctg gtgacagcaa ctatccctac tcctactacg aaggcaacga  540
    13031 ctttgcactg aaccttgcca tcccggacat tgactttggc actctccatc tctacaccac  600
W--> 13033 cgactgtacg tatctnccgg atgacccagt tctgcgttct ctctccatat cgggtgctaa  660
    13035 tgacttccag ggggcgtgag caacaactcg tggggcaaca agtgggtaca ag           712
    13632 <210> SEQ ID NO: 183
    13633 <211> LENGTH: 1245                                IMPORTANT
    13634 <212> TYPE: DNA                      The types of errors shown exist throughou
    13635 <213> ORGANISM: Aspergillus nidulans   insert <2207  the Sequence Listing. Please check subseq
    13637 <223> OTHER INFORMATION: unsure at all n locations   sequences for similar errors.
E--> 13638 <400> SEQUENCE: 183
    13640 atgataagca cagcgtatcc cgcttggggt aacgacggcg agggaatagc accggacttt   60
    13642 ataacatcgc tgtcaaaagc tccgacctct cattgagtat ccgtagtttg tctgcctcaa  120
    13644 agaaatagaa aagatttggg tgaccgtaag acaaacacac ggaatggcac aagcaaggcc  180
    13646 agcgagtgtc tctggaccga cagtgggagt gtcaagacag ggcgagacgc agcgtatgga  240
    13648 atgatcgttc ctagttgtcc ctgtaagaaa catcgaagtg gggtagtcga ttgcactaca  300
    13650 agtataacga tcttgagtgg ctgaacggag gaaaacgggc agaaccgctt cagtaacagc  360
    13652 tgagcctcgt gagtggtcga atctgcatct taatttttc tcagcacagt gacagtttct  420
    13654 ggttgggcga gctggcgcct cttaagtaga gggtccgtag gctccataga tcagactgcc  480
    13656 tccgggataa aaggtcacat ctcccttcaa gcagtcaatg tacttacggg aaacatggaa  540
    13658 acatgagtca acttgcagaa gtgggcatag ctttgttatc tccgttgcag gtcgggcact  600
    13660 tggtaagctc acttctttgt cgagtactga atatacccat atttcgaaga gaattatggt  660
    13662 tgtgctgttc gtgagttgtc ggatggctgc cgattgtaca tcaagaccgg tgcttgggat  720
    13664 tgccgaaatc cttgctgctg ccggtaagat gccagcgtct ggtatctccc tggtcaagga  780
    13666 ctcttgggca acttaagacc tattttcaat catgaaatca gcgttcattg agtgagtgta  840
    13668 gagtgatttc gaagaagtat acttgccgac ggttggaaaa acaggaacgc tggttccaca  900
    13670 ccaaatggct gaccgccgac cattccctac tcggaagagg aatcacgagt ctgctgatgt  960
```

RAW SEQUENCE LISTING   DATE: 01/07/2004
PATENT APPLICATION: US/09/404,520A   TIME: 12:07:42

                Input Set : **E:\ENU.corrected SEQ list 12.23.03.rpt.txt**
                Output Set: **N:\CRF4\01072004\I404520A.raw**

```
         13672 catcggttcc acgtgatgca cgcgcggtct tgagcttccc caagctgggt actgggagaa 1020
         13674 gcttcaggtt tcagcattaa gggtcttgtg caatgtcact gacgcccggt tagcgtccta 1080
W-->     13676 tcaatctcaa tctagctttc agcgtcttnt caagtccatc gccggtcctg tttcggatca 1140
         13678 ccatcgctat ctnccgtgct atcacggttt tccggcgtcc caccgagctt tcgctcctta 1200
         13680 gccgccctgg tccctccttc agctcagttt ctgtctgacc tagcg                1245
```

Important The types of errors shown exist throughout
the Sequence Listing. Please check subsequent
sequences for similar errors.

<210>   44347 *last sequence in submitted file*
<211>   21
<212>   DNA
<213>   Aspergillus nidulans

<400>   44347

gttctcgaga cgagcgatcg c   21

VARIABLE LOCATION SUMMARY   DATE: 01/07/2004
PATENT APPLICATION: US/09/404,520A   TIME: 12:08:40

Input Set : E:\ENU.corrected SEQ list 12.23.03.rpt.txt
Output Set: N:\CRF4\01072004\I404520A.raw

Use of n's or Xaa's(NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
in <220> to <223> section, please explain location of n or Xaa, and which
residue n or Xaa represents.

```
Seq#:24;  N Pos. 211
Seq#:28;  N Pos. 664
Seq#:33;  N Pos. 1020
Seq#:37;  N Pos. 1168
Seq#:45;  N Pos. 586,757,758
Seq#:47;  N Pos. 6706
Seq#:51;  N Pos. 274
Seq#:80;  N Pos. 3456,3461
Seq#:84;  N Pos. 619,624
Seq#:108; N Pos. 188
Seq#:117; N Pos. 750
Seq#:120; N Pos. 75
Seq#:129; N Pos. 656
Seq#:141; N Pos. 36,37,38,39,40,41,42,43,44,45,46,47,48,49,50,51,52,53,54
Seq#:141; N Pos. 55,56,57,58,59,60,61,62,63,64,65,66,67,68,69,70,71,72,73
Seq#:141; N Pos. 74,75,76,77,78,79,80,81,82,83,84,85,86,87
Seq#:144; N Pos. 1431
Seq#:149; N Pos. 351
Seq#:154; N Pos. 2054,2056
Seq#:166; N Pos. 86
Seq#:177; N Pos. 616
Seq#:183; N Pos. 1109,1153
Seq#:195; N Pos. 4195
Seq#:197; N Pos. 830
Seq#:205; N Pos. 3103
Seq#:209; N Pos. 785,793,824
Seq#:220; N Pos. 16,33,34,74
Seq#:233; N Pos. 447
Seq#:236; N Pos. 146
Seq#:245; N Pos. 316,411
Seq#:275; N Pos. 502
Seq#:285; N Pos. 1544
Seq#:304; N Pos. 912
Seq#:310; N Pos. 671
Seq#:324; N Pos. 1508
Seq#:325; N Pos. 1222,1223,2378,2382,2427,2608
Seq#:334; N Pos. 803,807
Seq#:347; N Pos. 2863
Seq#:355; N Pos. 5912
Seq#:360; N Pos. 2289
Seq#:363; N Pos. 18
Seq#:365; N Pos. 519
Seq#:367; N Pos. 2039
```

VARIABLE LOCATION SUMMARY   DATE: 01/07/2004
PATENT APPLICATION: **US/09/404,520A**   TIME: 12:08:40

Input Set : **E:\ENU.corrected SEQ list 12.23.03.rpt.txt**
Output Set: **N:\CRF4\01072004\I404520A.raw**

Seq#:403; N Pos. 73
Seq#:425; N Pos. 620
Seq#:426; N Pos. 1004
Seq#:439; N Pos. 47,50,68,138,162,165,243,250,264,272,298,299,301,302,334
Seq#:439; N Pos. 385,420,463,489,493,573
Seq#:441; N Pos. 423
Seq#:457; N Pos. 669
Seq#:460; N Pos. 4388,4397

Input Set : **E:\ENU.corrected SEQ list 12.23.03.rpt.txt**
Output Set: **N:\CRF4\01072004\I404520A.raw**

L:13 M:270 C: Current Application Number differs, Replaced Current Application No
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:1363 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:24
L:1363 M:283 W: Missing Blank Line separator, <400> field identifier
L:1371 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24
L:1371 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:24
L:1371 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:24
L:1371 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:180
L:1560 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:28
L:1560 M:283 W: Missing Blank Line separator, <400> field identifier
L:1584 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:28
L:1584 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:28
L:1584 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:28
L:1584 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:660
L:1925 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:33
L:1925 M:283 W: Missing Blank Line separator, <400> field identifier
L:1959 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:33
L:1959 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:33
L:1959 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:33
L:1959 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:960
L:2066 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:37
L:2066 M:283 W: Missing Blank Line separator, <400> field identifier
L:2106 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:37
L:2106 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:37
L:2106 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:37
L:2106 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:1140
L:2411 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:45
L:2411 M:283 W: Missing Blank Line separator, <400> field identifier
L:2431 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:45
L:2431 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:45
L:2431 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:45
L:2431 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45 after pos.:540
M:341 Repeated in SeqNo=45
L:2468 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:47
L:2468 M:283 W: Missing Blank Line separator, <400> field identifier
L:2692 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:47
L:2692 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:47
L:2692 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:47
L:2692 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:47 after pos.:6660
L:2783 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:51
L:2783 M:283 W: Missing Blank Line separator, <400> field identifier
L:2793 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:51
L:2793 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:51
L:2793 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:51
L:2793 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51 after pos.:240
L:7706 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:80
L:7706 M:283 W: Missing Blank Line separator, <400> field identifier
L:7822 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:80

Input Set : **E:\ENU.corrected SEQ list 12.23.03.rpt.txt**
Output Set: **N:\CRF4\01072004\I404520A.raw**

L:7822 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:80
L:7822 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:80
L:7822 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:80 after pos.:3420
L:8087 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:84
L:8087 M:283 W: Missing Blank Line separator, <400> field identifier
L:8109 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:84
L:8109 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:84
L:8109 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:84
L:8109 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:84 after pos.:600
L:9224 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:108
L:9224 M:283 W: Missing Blank Line separator, <400> field identifier
L:9232 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:108
L:9232 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:108
L:9232 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:108
L:9232 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:108 after pos.:180
L:9721 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:117
L:9721 M:283 W: Missing Blank Line separator, <400> field identifier
L:9747 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:117
L:9747 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:117
L:9747 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:117
L:9747 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:117 after pos.:720
L:9832 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:120
L:9832 M:283 W: Missing Blank Line separator, <400> field identifier
L:9836 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:120
L:9836 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:120
L:9836 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:120
L:9836 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:120 after pos.:60
L:10081 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:129
L:10081 M:283 W: Missing Blank Line separator, <400> field identifier
L:10103 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:129
L:10103 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:129
L:10103 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:129
L:10103 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:129 after pos.:600
L:10596 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:141
L:10596 M:283 W: Missing Blank Line separator, <400> field identifier
L:10598 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:141
L:10598 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:141
L:10598 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:141
L:10598 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:141 after pos.:0
M:341 Repeated in SeqNo=141
L:10791 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:144
L:10791 M:283 W: Missing Blank Line separator, <400> field identifier
L:10839 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:144
L:10839 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:144
L:10839 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:144
L:10839 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:144 after pos.:1380
L:11010 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:149
L:11010 M:283 W: Missing Blank Line separator, <400> field identifier
L:11022 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:149

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L:11022 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:149
L:11022 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:149
L:11022 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:149 after pos.:300
L:11285 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:154
L:11285 M:283 W: Missing Blank Line separator, <400> field identifier
L:11355 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:154
L:11355 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:154
L:11355 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:154 after pos.:2040
L:11964 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:166
L:11964 M:283 W: Missing Blank Line separator, <400> field identifier
L:11968 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:166 after pos.:60
L:13011 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:177
L:13011 M:283 W: Missing Blank Line separator, <400> field identifier
L:13033 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:177 after pos.:600
L:13638 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:183
L:13638 M:283 W: Missing Blank Line separator, <400> field identifier
L:13676 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:183 after pos.:1080
M:341 Repeated in SeqNo=183
L:14911 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:195
L:14911 M:283 W: Missing Blank Line separator, <400> field identifier
L:15051 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:195 after pos.:4140
L:15118 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:197
L:15118 M:283 W: Missing Blank Line separator, <400> field identifier
L:15146 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:197 after pos.:780
L:15673 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:205
L:15673 M:283 W: Missing Blank Line separator, <400> field identifier
L:15777 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:205 after pos.:3060
L:16028 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:209
L:16028 M:283 W: Missing Blank Line separator, <400> field identifier
L:16056 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:209 after pos.:780
L:16661 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:220
L:16661 M:283 W: Missing Blank Line separator, <400> field identifier
L:16663 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:220 after pos.:0
M:341 Repeated in SeqNo=220
L:17374 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:233
L:17374 M:283 W: Missing Blank Line separator, <400> field identifier
L:17390 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:233 after pos.:420
L:17457 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:236
L:17457 M:283 W: Missing Blank Line separator, <400> field identifier
L:17463 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:236 after pos.:120
L:17884 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:245
L:17884 M:283 W: Missing Blank Line separator, <400> field identifier
L:17896 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:245 after pos.:300
M:341 Repeated in SeqNo=245
L:19129 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:275
L:19129 M:283 W: Missing Blank Line separator, <400> field identifier
L:19147 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:275 after pos.:480
L:19734 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:285
L:19734 M:283 W: Missing Blank Line separator, <400> field identifier
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VERIFICATION SUMMARY   DATE: 01/07/2004
PATENT APPLICATION: **US/09/404,520A**   TIME: 12:08:40

Input Set : **E:\ENU.corrected SEQ list 12.23.03.rpt.txt**
Output Set: **N:\CRF4\01072004\I404520A.raw**

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L:19786 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:285 after pos.:1500
L:20987 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:304
L:20987 M:283 W: Missing Blank Line separator, <400> field identifier
L:21019 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:304 after pos.:900
L:21306 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:310
L:21306 M:283 W: Missing Blank Line separator, <400> field identifier
L:21330 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:310 after pos.:660
L:21915 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:324
L:21915 M:283 W: Missing Blank Line separator, <400> field identifier
L:21967 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:324 after pos.:1500
L:21988 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:325
L:21988 M:283 W: Missing Blank Line separator, <400> field identifier
L:22030 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:325 after pos.:1200
M:341 Repeated in SeqNo=325
L:22577 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:334
L:22577 M:283 W: Missing Blank Line separator, <400> field identifier
L:22605 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:334 after pos.:780
L:23450 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:347
L:23450 M:283 W: Missing Blank Line separator, <400> field identifier
L:23546 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:347 after pos.:2820
L:24201 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:355
L:24201 M:283 W: Missing Blank Line separator, <400> field identifier
L:24399 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:355 after pos.:5880
L:24620 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:360
L:24620 M:283 W: Missing Blank Line separator, <400> field identifier
L:24698 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:360 after pos.:2280
L:24853 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:363
L:24853 M:283 W: Missing Blank Line separator, <400> field identifier
L:24855 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:363 after pos.:0
L:25104 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:365
L:25104 M:283 W: Missing Blank Line separator, <400> field identifier
L:25122 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:365 after pos.:480
L:25187 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:367
L:25187 M:283 W: Missing Blank Line separator, <400> field identifier
L:25255 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:367 after pos.:1980
L:28090 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:403
L:28090 M:283 W: Missing Blank Line separator, <400> field identifier
L:28094 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:403 after pos.:60
L:30085 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:425
L:30085 M:283 W: Missing Blank Line separator, <400> field identifier
L:30107 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:425 after pos.:600
L:30118 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:426
L:30118 M:283 W: Missing Blank Line separator, <400> field identifier
L:30152 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:426 after pos.:960
L:30961 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:439
L:30961 M:283 W: Missing Blank Line separator, <400> field identifier
L:31028 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:441
L:31028 M:283 W: Missing Blank Line separator, <400> field identifier
L:32189 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:457
```

Input Set : **E:\ENU.corrected SEQ list 12.23.03.rpt.txt**
Output Set: **N:\CRF4\01072004\I404520A.raw**

L:32189 M:283 W: Missing Blank Line separator, <400> field identifier
L:32294 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:460
L:32294 M:283 W: Missing Blank Line separator, <400> field identifier
L:32535 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:463
L:32535 M:283 W: Missing Blank Line separator, <400> field identifier
L:32598 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:464
L:32598 M:283 W: Missing Blank Line separator, <400> field identifier
L:13 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (44345) Counted (44347)